

Preliminary Amendment

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Applicant(s): Benson et al.

Serial No.: 10/028,224

Confirmation No.: 4497

Filed: December 21, 2001

*For: CRYSTALLIZATION AND STRUCTURE DETERMINATION OF GLYCOSYLATED HUMAN BETA
SECRETASE, AN ENZYME IMPLICATED IN ALZHEIMER'S DISEASE*

Remarks

The Examiner alleged that the present application failed to comply with the requirements of 37 C.F.R. §§1.821-1.825. The specification has been amended to recite a SEQ ID NO. along with each sequence in the specification, in accordance with 37 C.F.R. §1.821 through §1.825. The amendment is supported, for example, by the Sequence Listing submitted by Applicants on December 21, 2001.

Since the SEQ ID NOs are already included in the paper copy and computer readable form of the Sequence Listing submitted by Applicants on December 21, 2001, Applicants respectfully submit that no further correction is required.

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Conclusion

The Examiner is invited to contact Applicants' Representatives at the below-listed telephone number, if there are any questions regarding this Response to Restriction Requirement and Preliminary Amendment or if prosecution of this application may be assisted thereby.

Respectfully submitted for
Benson et al.

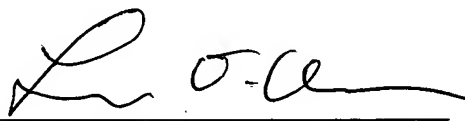
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PATENT TRADEMARK OFFICE


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I hereby certify that this paper and/or fee is/are being deposited with the United States Postal Service "Express Mail Post Office to Addressee" service under 37 C.F.R. 1.10 on the date indicated above and is addressed to the Assistant Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450.


Name: **SAM HER**



APPENDIX A - SPECIFICATION/CLAIM AMENDMENTS
INCLUDING NOTATIONS TO INDICATE CHANGES MADE

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Docket No.: 00403.CN1

Amendments to the following are indicated by underlining what has been added and bracketing what has been deleted. Additionally, all amendments have been indicated by the use of bold font.

In the Specification

The paragraph beginning at page 12, line 2, has been amended as follows:

Figure 2 is the synthetic peptide Ser-Glu-Val-Asn-Sta-Val-Ala-Glu-Phe-Arg-Gly-Gly-Cys (where Sta = statine) **(SEQ ID NO:3)**, PNU-292593E, used for affinity purification of beta secretase.

The paragraph beginning at page 40, line 25, has been amended as follows:

The expression plasmid = pcDNA3.1/myc/his (neomycin) (Invitrogen) contains beta secretase extending from Met [-21] to Ser [432] with a myc tag followed by a hexahistidine tag [EQKLISEEDLN**MHTEHHHHHH***] **(SEQ ID NO:2)** at the C-terminus. Following transfection in HEK293 cells, stable cells were selected using 0.8 mg/ml G418. A stable clone of transfected HEK293 cells that secretes human beta-secretase was expanded in static, monolayer cell culture. Confluent cultures were detached by shaking and a plurality of plastic, 225 cm² T-flasks were each inoculated with a suspension of 1-5 x 10⁶ cells in 100 ml of High-Glucose Dulbecco's Modified Eagle medium that was supplemented with 5% fetal bovine serum and 500 micrograms/ml G418. These cell cultures were incubated in a humidified, 37°C incubator gassed with 95% air and 5% CO₂. Once the cells reached confluence the growth medium in each flask was removed and replaced with 100 ml fresh medium. The conditioned, culture medium supernatant was harvested aseptically and replaced by fresh medium every 48-72 hours. The harvested medium was pooled, centrifuged at 1000 x g to remove cell debris, and was stored in plastic bottles at 4°C. Cell monolayers were maintained in semi-continuous culture for several weeks until the cells either began to die or to detach from the culture flasks. The cells were then resuspended and used to inoculate a fresh set of production flasks.

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The paragraph beginning at page 42, line 20, has been amended as follows:

Production of Recombinant Human β -Secretase in Insect sf9 Cells and CHO-K1 Cells.

The coding sequence was engineered to delete the terminal transmembrane and cytoplasmic domain and introduce a C-terminal hexahistidine tag using the polymerase chain reaction. The 5' sense oligonucleotide primer [CGCTTTGGATCCGTGGACAACCTGAGGGGCAA] (**SEQ ID NO:4**) was designed to incorporate a BamHI site for ease in subcloning and Kozak consensus sequence around the initiator methionine for optimal translation initiation. The 3' antisense primer [CGCTTTGGTACCCTATGACTCATCTGTCTGTGGAATGTTG] (**SEQ ID NO:5**) incorporated a hexahistidine tag and translation termination codon just upstream of the predicted transmembrane domain (Ser⁴³²) and a NotI restriction site for cloning. The PCR was performed on the plasmid template pcDNA3.1hygroAsp2R for 15 cycles [94°C, 30 sec., 65°C, 30 sec., 72°C, 30 sec] using Pwo I polymerase (Roche Biochemicals, Indianapolis, IN) as outlined by the manufacturer. The PCR product was digested to completion with *Bam*HI and *Not*I and ligated into the *Bam*HI and *Not*I sites of the Baculovirus transfer vector pVL1393 (PharMingen, San Diego, CA). A portion of the ligation was used to transform competent E. coli DH5 α cells and recombinant clones were selected on ampicillin. Individual clones containing the proper cDNA inserts were identified by PCR. Plasmid DNA from clone (pVL1393/Hu_Asp-2L Δ TM(His)₆) was prepared by alkaline lysis and banding in CsCl. The integrity of the insert was confirmed by complete DNA sequencing. For CHO-K1 cell expression, plasmid pVL1393/Hu_Asp-2L Δ TM(His)₆ was digested with *Bam*HI and *Not*I and the resulting fragment subcloned into the mammalian expression vector pcDNA3.1(hygro) as described above to yield pcDNA3.1(hygro)/Hu_Asp-2L Δ TM(His)₆.

The paragraph beginning at page 44, line 28, has been amended as follows:

The 40-80% ammonium sulfate pellet was dissolved in 25 mM Tris-HCl (8.5)/0.5 M NaCl/10 mM imidazole (1/10 the original volume) and applied to a 12.5 ml column containing Ni⁺-NTA Fast Flow resin previously equilibrated in the same buffer. Following sample

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application, the column was washed with 10 column volumes of loading buffer and then eluted with 25 mM Tris-HCl (8.5)/0.5 M NaCl/50 mM imidazole. The material eluting in 50 mM imidazole was pooled, concentrated approximately 10-fold using a YM 30 membrane (30,000 MWCO), and then dialyzed against 10 mM HEPES-Na (8.0) using 50,000 molecular weight cutoff tubing. For affinity purification, the synthetic peptide Ser-Glu-Val-Asn-Sta-Val-Ala-Glu-Phe-Arg-Gly-Gly-Cys (where Sta = statine, PNU-292593E) (**SEQ ID NO:3**) was synthesized and coupled to sulfolink resin (Pierce Chemical Company) as recommended by the manufacture. The dialyzed material from above was adjusted to 0.1 M NaOAc (4.5) by addition of 1/10 volume of 1.0 M NaOAc (4.5) and immediately applied to the PNU-292593E/sulfolink column (6 ml containing 1.0 mg PNU-292593/ml of resin) that had been previously equilibrated in 25 mM NaOAc (4.5). Following sample application, the column was washed with 10 column volumes of 25 mM NaOAc (4.5) and then eluted with 50 mM NaBO₃ (8.5). N-terminal sequence analysis of the affinity purified material revealed an equimolar mixture of pro- and processed human β -secretase beginning at Thr¹ and Glu²⁵, respectively. The final protein concentration was determined by amino acid analysis assuming a 52 kDa glycoprotein for insect cells and a 60 kDa glycoprotein for CHO cells, respectively.